
Sequence Listing was accepted with existing errors.

See attached Validation Report.

If you need help call the Patent Electronic Business Center at (866)

217-9197 (toll free).

Reviewer: Anne Corrigan

Timestamp: Thu Jun 07 20:32:02 EDT 2007

Validated By CRFValidator v 1.0.2

Application No: 10633850 Version No: 2.1

Input Set:

Output Set:

Started: 2007-06-07 20:31:45.592 **Finished:** 2007-06-07 20:31:49.766

Elapsed: 0 hr(s) 0 min(s) 4 sec(s) 174 ms

Total Warnings: 29

No. of SeqIDs Defined: 113

Actual SeqID Count: 113

Total Errors:

Error code		Error Description								
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Input Set:

Output Set:

Started: 2007-06-07 20:31:45.592

Finished: 2007-06-07 20:31:49.766

Elapsed: 0 hr(s) 0 min(s) 4 sec(s) 174 ms

Total Warnings: 29

Total Errors: 1

No. of SeqIDs Defined: 113

Actual SeqID Count: 113

Error code Error Description

W 213 Artificial or Unknown found in <213> in SEQ ID (45)

This error has occured more than 20 times, will not be displayed

SEQUENCE LISTING

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Leu	Thi	Tyr	Pro 20	Gln	Cys	Ser	Leu	Ser 25	Lys	Glu	Glu	Ser	Leu 30	Ser	Gln
Leu	Glr	n Ala 35	Leu	Asn	Thr	Pro	Ile 40	Asn	Lys	Lys	Phe	Ile 45	Lys	Ile	Суз
Arg	Glu 50	ı Leu	His	Glu	Asp	Gly 55	Gln	Pro	His	Leu	His	Val	Leu	Ile	Gln
Phe 65	Glu	ı Gly	Lys	Tyr	Cys 70	Суз	Gln	Asn	Gln	Arg 75	Phe	Phe	Asp	Leu	Val 80
Ser	Pro) Thr	Arg	Ser 85	Ala	His	Phe	His	Pro 90	Asn	Ile	Gln	Arg	Ala 95	Lys

Ser Ser Ser Asp Val Lys Thr Tyr Ile Asp Lys Asp Gly Asp Thr Leu
100 105 110

Val '	Trp	Gly 115	Glu	Phe	Gln	Val	Asp 120	Gly	Arg	Ser	Ala	Arg 125	Gly	Gly	Суз
Gln '	Thr 130	Ser	Asn	Asp	Ala	Ala 135	Ala	Glu	Ala	Leu	Asn 140	Ala	Ser	Ser	Lys
Glu (Glu	Ala	Leu	Gln	Ile 150	Ile	Arg	Glu	Lys	Ile 155	Pro	Glu	Lys	Tyr	Leu 160
Phe (Gln	Phe	His	Asn 165	Leu	Asn	Ser	Asn	Leu 170	Asp	Arg	Ile	Phe	Asp 175	Lys
Thr 1	Pro	Glu	Pro 180	Trp	Leu	Pro	Pro	Phe 185	His	Val	Ser	Ser	Phe 190	Thr	Asn
Val 1	Pro	Asp 195	Glu	Met	Arg	Gln	Trp 200	Ala	Glu	Asn	Tyr	Phe 205	Gly	Lys	Ser
Ser .	Ala 210	Ala	Arg	Pro	Glu	Arg 215	Pro	Ile	Ser	Ile	Ile 220	Ile	Glu	Gly	Asp
Ser 2	Arg	Thr	Gly	Lys	Thr 230	Met	Trp	Ala	Arg	Ser 235	Leu	Gly	Pro	His	Asn 240
Tyr :	Leu	Ser	Gly	His 245	Leu	Asp	Leu	Asn	Ser 250	Arg	Val	Tyr	Ser	Asn 255	Lys
Val (Glu	Tyr	Asn 260	Val	Ile	Asp	Asp	Val 265	Thr	Pro	Gln	Tyr	Leu 270	Lys	Leu
Lys	His	Trp 275	Lys	Glu	Leu	Ile	Gly 280	Ala	Gln	Arg	Asp	Trp 285	Gln	Thr	Asn
Cys :	Lys 290	Tyr	Gly	Lys	Pro	Val 295	Gln	Ile	Lys	Gly	Gly 300	Ile	Pro	Ser	Ile
Val :	Leu	Cys	Asn	Pro	Gly 310	Glu	Gly	Ala	Ser	Tyr 315	Lys	Val	Phe	Leu	Asp 320
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                                 45
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Ser Lys Glu Glu Ala Leu Gln Ile Ile Arg Glu Lys Ile Pro Glu Lys
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                       25
         20
                                     30
Ser Ala Ala Ala Leu Gln Ile Ile Arg Glu Lys Ile Pro Glu Lys
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Tyr Leu Phe Gln Phe His Asn Leu Asn Ser Asn Leu Asp Arg Ile Phe
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Ser Lys Glu Glu Ala Leu Gln Ile Ile Ala Ala Ile Pro Glu Lys 35 40 45

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Asp Lys Thr Pro Glu Pro 65 70

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Tyr Leu Phe Ala Phe Ala Ala Leu Asn Ser Asn Leu Asp Arg Ile Phe
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       20
               25
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Ser Lys Glu Glu Ala Leu Gln Ile Ile Arg Glu Lys Ile Pro Glu Lys
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 35
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                       10
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                25
        20
Ser Lys Glu Glu Ala Leu Gln Ile Ile Arg Glu Lys Ile Pro Glu Lys
     35
                 40
                                      45
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Asp Lys Thr Pro Glu Pro
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1 5
Gly Cys Gln Thr Ser Asn Asp Ala Ala Ala Glu Ala Leu Asn Ala Ser
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         20
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Ser Ala Ala Ala Leu Gln Ile Ile Arg Glu Lys Ile Pro Glu Lys 35 40 45

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                   25
Ser Lys Glu Glu Ala Leu Gln Ile Ile Arg Glu Lys Ile Pro Glu Lys
       35
                40
Tyr Leu Phe Gln Phe His Asn Leu Asn Ser Asn Leu Asp Arg Ile Phe
                     55
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Asp Ala Thr Pro Ala Pro
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Asp Lys Thr Pro Glu Pro 65 70

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Tyr Leu Phe Gln Phe His Asn Leu Asn Ser Asn Leu Asp Arg Ile Phe
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Asp Lys Thr Pro Glu Pro
65 70
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         20
                                        30
Ser Lys Ala Ala Ala Leu Gln Ile Ile Arg Glu Lys Ile Pro Glu Lys
   35 40 45
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